

**Protein sequences of th examples (mutated to yield soluble forms)**

&gt;sFcγRI

SEQ ID NO: 1

MAVISLQPPWVSVFQEETVTLHCEVLHLPGSSSTQWFLNGTATQTSTPSYRITSASVND  
GEYRCQRGLSGRSDPIQLEIHRGWLLQVSSRVFTEGEPLALRCHAWKDKLVYNVLYRN  
GKAFKFFHWNNSNLTKTNISHNGTYHCSGMGKHRYTSAGISVTVKELFPAPVLNASVTS  
PLLEGNLVTLSKETKLLQRPGLQLYFSFYMGSKTLRGRNTSSEYQILTARREDSGLYWC  
EAATEDGNVLKRSPELELQVLGLQLPTPV

SEQ ID NO: 2

&gt;sFcγRIIa

MAAPPKAVLKLEPPWINVLQEDSVTLTCQGARSPESDSIQWFHNGNLIPTHTQPSYRFA  
NNNDSGEYTCQTGQTSLSDPVHLTVLSEWLVLQTPHLEFQEGETIMLRCHSWKDKPLVKV  
TFFQNGKSQKFSRLDPTFSIPQANHSHSGDYHCTGNIGYTLFSSKPVITITVQVP

SEQ ID NO: 3

&gt;sFcγRIIb

AGTPAAPKAVLKLEPPWINVLQEDSVTLTCRGTHSPESDSIQWFHNGNLIPTHTQPSYR  
FKANNDSGEYTCQTGQTSLSDPVHLTVLSEWLVLQTPHLEFQEGETIVLRCHSWKDKPL  
VKVTFQNGKSKKFSRSDPNFSIPQANHSHSGDYHCTGNIGYTLYSSKPVITITVQAPSS  
PMGII

SEQ ID NO: 4

&gt;sFcγRIII

MRTEDLPKAVVFLEPQWYSVLEKDSVTLKCGAYSPEDNSTQWFHNESLISSQASSYFID  
AATVNDSEYRCQTNLSTLSDPVQLEVHIGWLLQAPRWVFKEEDPIHLRCHSWKNTALH  
KVTYLQNGKDRKYFHNSDFHIPKATLKDSGSYFCRGLVGSKNVSSETVNITITQG

SEQ ID NO: 5

&gt;sFcεRIa

MAVPQKPKVSLNPPWNRIFKGENVTLCNGNFFEVSSTKWFHNGSLSEETNSSLNIVN  
AKFEDSGEYKQHQVNESEPVYLEVFSDWLLQASAEVVMGQPLFLRCHGWRNWDVYK  
VIYYKDGEALKYWYENHNISITNATVEDSGTYCTGKVVQLDYESEPLNITVIKAPREKY  
WLQF

SEQ ID NO: 6

&gt;sFcεRII

MDTTQSLKQLEERAARNVSQVSKNLESHHGDMTQKSQSTQISQELEELRAEQRLKSQD  
LELSWNLNGLQADLSSFKSQELNERNEASDLLERLREEVTKLRMELQVSSGFVCNTCPEK  
WINFQRKCYFFGKGTKQWVHARYACDDMEGQLVSIHSPEEQDFLTKHASHTGSWIGLRNL  
DLKGEFIWVDGSHVDYSNAPGEPTSRSQGEDCVMRGRSGRWNDADFCDRKLGAWVCDRLA  
TCTPPASEGSAESMGPDSPDPDGRLLPTPSAPLHS

09856933-022702

**DNA sequences of the examples (mutated to yield soluble forms)**

		SEQ ID NO: 7						
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1	CATATGGCAG	TGATCTCTTT	GCAGCCTCCA	TGGGTACGCG	TGTTCCAAGA	GGAAACCGTA	60	
61	ACCTTGCACT	GTGAGGTGCT	CCATCTGCCT	GGGAGCAGCT	CTACACAGTG	GTTTCTCAAT	120	
121	GGCACAGCCA	CTCAGACCTC	GACCCCCAGC	TACAGAATCA	CCTCTGCCAG	TGTCAATGAC	180	
181	AGTGGTGAAT	ACAGGTGCCA	GAGAGGTCTC	TCAGGGCGAA	GTGACCCCAT	ACAGCTGGAA	240	
241	ATCCACAGAG	GCTGGCTACT	ACTGCAGGTC	TCCAGCAGAG	TCTTCACGGA	AGGAGAACCT	300	
301	CTGGCCTTGA	GGTGTCTATG	GTGGAAGGAT	AAGCTGGTGT	ACAATGTGCT	TTACTATCGA	360	
361	AATGGCAAAG	CCTTTAAGTT	TTTCCACTGG	AATTCTAACC	TCACCATTCT	GAAAACCAAC	420	
421	ATAAGTCACA	ATGGCACCTA	CCATTGTCTA	GGCATGGGAA	AGCATCGCTA	CACATCAGCA	480	
481	GGAATATCTG	TCACTGTGAA	AGAGCTATTT	CCAGCTCCAG	TGCTGAATGC	ATCTGTGACA	540	
541	TCCCCACTCC	TGGAGGGGAA	TCTGGTCAAC	CTGAGCTGTG	AAACAAAGTT	GCTCTTGCAG	600	
601	AGGCCTGGTT	TGCAGCTTTA	CTTCTCCTTC	TACATGGGCA	GCAAGACCCT	GCGAGGCAGG	660	
661	AACACATCCT	CTGAATACCA	AATACTAACT	GCTAGAAGAG	AAGACTCTGG	GTTATACTGG	720	
721	TGCGAGGCTG	CCACAGAGGA	TGGAAATGTC	CTTAAGCGCA	GCCCTGAGTT	GGAGCTTCAA	780	
781	GTGCTTGGCC	TCAGTTACC	AACTCCTGTC	TAGTCTCGAG			820	
		SEQ ID NO: 8						
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1	CATATGGCAG	CTCCCCAAA	GGCTGTGCTG	AAACTTGAGC	CCCCGTGGAT	CAACGTGCTC	60	
61	CAGGAGGACT	CTGTGACTCT	GACATGCCAG	GGGGCTCGCA	GCCCTGAGAG	CGACTCCATT	120	
121	CAGTGGTTCC	ACAATGGGAA	TGTCATTCCC	ACCCACACGC	AGCCCAGCTA	CAGGTTCAAG	180	
181	GCCAACAACA	ATGACAGCGG	GGAGTACACG	TGCCAGACTG	GCCAGACCAG	CCTCAGCGAC	240	
241	CCTGTGCATC	TGACTGTGCT	TTCCGAATGG	CTGGTGCTCC	AGACCCCTCA	CCTGGAGTTC	300	
301	CAGGAGGGAG	AAACCATCAT	GCTGAGGTGC	CACAGCTGGA	AGGACAAGCC	TCTGGTCAAG	360	
361	GTCACATTCT	TCCAGAATGG	AAAATCCCAG	AAATTCTCCC	GTTTGGATCC	CACCTTCTCC	420	
421	ATCCACAAG	CAAACCACAG	TCACAGTGGT	GATTACCACT	GCACAGGAAA	CATAGGCTAC	480	
481	ACGCTGTTCT	CATCCAAGCC	TGTGACCATC	ACTGTCCAAG	TGCCCTGAAG	CTT	533	
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61	ACGTGCTCCA	GGAGGACTCT	GATGACTCTGA	CATGCCGGGG	GACTCACAGC	CCTGAGAGCG	120	
121	ACTCCATTCA	GTGGTTCCAC	AATGGGAATC	TCTATCCCAC	CCACACGCAG	CCCAGCTACA	180	
181	GGTTCAAGGC	CAACAACAAT	GACAGCGGGG	AGTACACGTG	CCAGACTGGC	CAGACCAGCC	240	
241	TCAGCGACCC	TGTGCATCTG	ACTGTGCTTT	CTGAGTGGCT	GGTGCTCCAG	ACCCCTCACC	300	
301	TGGAGTTCCA	GGAGGGAGAA	ACCATCGTGC	TGAGGTGCCA	CAGCTGGAAG	GACAAGCCTC	360	
361	TGGTCAAGGT	CACATTCTTC	CAGAATGGAA	AATCCAAGAA	ATTTTCCCGT	TCGGATCCCA	420	
421	ACTTCTCCAT	CCCACAAGCA	AACCACAGTC	ACAGTGGTGA	TTACCACTGC	ACAGGAAACA	480	
481	TAGGCTACAC	GCTGTACTCA	TCCAAGCCTG	TGACCATCAC	TGTCCAAGCT	CCCAGCTCTT	540	
541	CACCGATGGG	GATCATTTAG	GCTGTCTGAC				569	
		SEQ ID NO: 10						
>sFcyRIII								
1	CATATGCGGA	CTGAAGATCT	CCCCAAGGCT	GTGGTGTTC	TGGAGCCTCA	ATGGTACAGC	60	
61	GTGCTTGAGA	AGGACAGTGT	GACTCTGAAG	TGCCAGGGAG	CCTACTCCCC	TGAGGACAAT	120	
121	TCCACACAGT	GGTTTCACAA	TGAGAGCCTC	ATCTCAAGCC	AGGCCTCGAG	CTACTTCATT	180	
181	GACGCTGCCA	CAGTCAACGA	CAGTGGAGAG	TACAGGTGCC	AGACAAACCT	CTCCACCCCTC	240	
241	AGTGACCCGG	TGCAGCTAGA	AGTCCATATC	GGCTGGCTGT	TGCTCCAGGC	CCCTCGGTGG	300	
301	GTGTTCAAGG	AGGAAGACCC	TATTCACCTG	AGGTGTCACA	GCTGGAAGAA	CACCTGCTCTG	360	
361	CATAAGGTCA	CATATTTACA	GAATGGCAAA	GACAGGAAGT	ATTTTCATCA	TAATTCTGAC	420	
421	TTCCACATTC	CAAAAGCCAC	ACTCAAAGAT	AGCGGCTCCT	ACTTCTGCAG	GGGGCTTGTT	480	
481	GGGAGTAAAA	ATGTGTCTTC	AGAGACTGTG	AACATCACCA	TCACTCAAGG	TTAAGCTT	538	
		SEQ ID NO: 11						
>sFceRIa								
1	CATATGGCAG	TCCCTCAGAA	ACCTAAGGTC	TCCTTGAACC	CTCCATGGAA	TAGAATATTT	60	
61	AAAGGAGAGA	ATGTGACTCT	TACATGTAAT	GGGAACAATT	TCTTTGAAGT	CAGTTCCACC	120	
121	AAATGGTTCC	ACAATGGCAG	CCTTTTCAGAA	GAGACAATT	CAAGTTTGAA	TATTGTGAAT	180	
181	GCCAAATTTG	AAGACAGTGG	AGAATACAAA	TGTCAGCACC	AACAAGTTAA	TGAGAGTGAA	240	
241	CCTGTGTACC	TGGAAGTCTT	CAGTGACTGG	CTGCTCCTTC	AGGCCTCTGC	TGAGGTGGTG	300	
301	ATGGAGGGCC	AGCCCTCTTT	CCTCAGGTGC	CATGGTTGGA	GGAAC TGGGA	TGTGTACAAG	360	
361	GTGATCTATT	ATAAGGATGG	TGAAGCTCTC	AAGTACTGGT	ATGAGAACCA	CAACATCTCC	420	
421	ATTACAAATG	CCACAGTTGA	AGACAGTGGA	ACCTACTACT	GTACGGGCAA	AGTGTGGCAG	480	
481	CTGGACTATG	AGTCTGAGCC	CCTCAACATT	ACTGTAATAA	AAGCTCCGCG	TGAGAAGTAC	540	
541	TGGCTACAAT	TTTAGGATCC					560	

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## SEQ ID NO: 12

&gt;sFceRII

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1 CATATGGAGT TGCAGGTGTC CAGCGGCTTT GTGTGCAACA CGTGCCCTGA AAAGTGGATC 60
61 AATTTCCAAC GGAAGTGCTA CTACTTCGGC AAGGGCACCA AGCAGTGGGT CCACGCCCGG 120
121 TATGCCCTGTG ACGACATGGA AGGGCAGCTG GTCAGCATCC ACAGCCCGGA GGAGCAGGAC 180
181 TTCCTGACCA AGCATGCCAG CCACACCGGC TCCTGGATTG GCCTTCGGAA CTTGGACCTG 240
241 AAGGGGGAGT TTATCTGGGT GGATGGGAGC CACGTGGACT ACAGCAACTG GGCTCCAGGG 300
301 GAGCCCACCA GCCGGAGCCA GGGCGAGGAC TCGGTGATGA TCGGGGGCTC CGGTCGCTGG 360
361 AACGACGCCT TCTGCGACCG TAAGCTGGGC GCCTGGGTGT GCGACCGGCT GGCCACATGC 420
421 ACGCCGCCAG CCAGCGAAGG TTCCGCGGAG TCCATGGGAC CTGATTCAAG ACCAGACCCT 480
481 GACGGCCGCC TGCCACCCCC CTCTGCCCCCT CTCCACTCTT GAGCATGGAT CC 532

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## SEQ ID NO:13

human FcγRIIb2

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1 ggctgtgact gctgtgctct gggcgccact cgctccaggg agtgatggga atcctgtcat
61 ttttacctgt ccttgccact gagagtgact gggctgactg caagtcccc cagccttggg
121 gtcatatgct tctgtggaca gctgtgctat tcttggtcc tggtgtggg acacctgcag
181 ccccccaaa ggctgtgctg aaactcgagc cccagtggat caactgtctc caggaggact
241 ctgtgactct gacatgccgg gggactcaca gccctgagag cgactccatt cagtggttcc
301 acaatgggaa tctcattccc acccacacgc agcccagcta caggttcaag gccaacaaca
361 atgacagcgg ggagtacacg tgccagactg gccagaccag cctcagcgac cctgtgcatc
421 tgacagtgtc ttctgagtgg ctggtgctcc agaccctca cctggagtgc caggaggag
481 aaaccatcgt gctgaggtgc cacagctgga aggacaagcc tctggtcaag gtcacattct
541 tccagaatgg aaaatccaag aaattttccc gttcggatcc caacttctcc atcccacaag
601 caaaccacag tcacagtggg gattaccatt gcacaggaaa cataggctac acgctgtact
661 catccaagcc tgtgaccatc actgtccaag ctcccagctc ttcaccgatg gggatcattg
721 tggctgtggg cactgggatt gctgtagctg ccattgttgc tgctgtagtg gccttgatct
781 actgcaggaa aaagcggatt tcagccaatc ccactaatcc tgatgaggct gacaaagtgt
841 gggctgagaa cacaatcacc tatteacttc tcatgcaccc ggatgctctg gaagagcctg
901 atgaccagaa ccgtatttag tctccattgt cttgcattgg gatttgagaa gaaatcagag
961 aggggaagatc tggattttcc tggcctaaat tccccttggg gaggacaggg agatgctgca
1021 gttccaaaag agaaggtttc ttccagagtc atctacctga gtccctgaagc tccctgtcct
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1201 ttattaacag ataatagcaa cttgggaaat gcttatgtta caggttacgt gagaacaatc
1261 atgtaaatct atatgatttc agaaatgtta aaatagacta acctctacca gcacattaaa
1321 agtgattggt tctgggtgat aaaattattg atgattttta ttttctttat tttctataa
1381 agatcatata ttacttttat aataaaacat tataaaaaac

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## SEQ ID NO:14

human FcεRIα

1 agatctcagc acagtaagca ccaggagtcc atgaagaaga tggctcctgc catggaatcc  
61 cctactctac tgtgtgtagc cttactgttc ttcgctccag atggcgtggt agcagtcctc  
121 cagaaacctc aggtctcctt gaacctcca tggaatagaa tatttaaagg agagaatgtg  
181 actcttacat gtaatgggaa caatttcttt gaagtcagtt ccaccaaatg gtccacaat  
241 ggcagccttt cagaagagac aaattcaagt ttgaatattg tgaatgccaa atttgaagac  
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361 gtcttcagtg actggctgct ccttcaggcc tctgctgagg tggatgagg gggccagccc  
421 ctcttcctca ggtgccatgg ttggaggaaac tgggatgtgt acaaggtgat ctattataag  
481 gatggtgaag ctctcaagta ctggtatgag aaccacaaca tctccattac aaatgccaca  
541 gttgaagaca gtggaacctc ctactgtacg ggcaaagtgt ggcagctgga ctatgagtct  
601 gagccccctc acattactgt aataaaagct ccgctgaga agtactggct acaatttttt  
661 atcccattgt tgggtggtgat tctgtttgct gtggacacag gattatttat ctcaactcag  
721 cagcagggtc catttctctt gaagattaag agaaccagga aaggcttcag acttctgaac  
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841 cattagtttt tttccagcat cagcaattgc tactcaattg tcaaacacag cttgcaatat  
901 acatagaaac gtctgtgctc aaggatttat agaaatgctt cattaaactg agtgaactg  
961 attaagtggc atgtaatagt aagtgtcaa ttaacattgg ttgaataaat gagagaatga  
1021 atagattcat ttattagcat ttgtaaaaga gatgttcaat ttagatct

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## SEQ ID NO:15

human mRNA for high affinity Fc receptor (FcγRI)

1 gacagatttc actgctccca ccagcttgga gacaacatgt ggttcttgac aactctgctc  
61 ctttgggttc cagttgatgg gcaagtggac accacaaagg cagtgatctc tttgcagcct  
121 ccattgggtca gcgtgttcca agaggaaacc gtaaccttgc actgtgaggt gctccatctg  
181 cctgggagca gctctacaca gtggtttctc aatggcacag ccaactcagac ctcgaccccc  
241 agctacagaa tcacctctgc cagtgtcaat gacagtgggtg aatacagggtg ccagagaggt  
301 ctctcagggc gaagtgaacc catacagctg gaaatccaca gaggtgggt actactgcag  
361 gtctccagca gagtcttcac ggaaggagaa cctctggcct tgagggtgtca tgcgtggaag  
421 gataagctgg tgtacaatgt gctttactat cgaaatggca aagcctttaa gtttttccac  
481 tgggaattcta acctcaccat tctgaaaacc aacataagtc acaatggcac ctaccattgc  
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1201 tgcccacttg cccccgtga gcaactgcgtg caaacatcca aaagttcaac aacaccagaa  
1261 ctgtgtgtct catggtatgt aactcttaaa gcaaataaat gaactgactt caaaaaaaaa  
1321 a

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## SEQ ID NO:16

human FcγRIIa

1 cccaaatgtc tcagaatgta tgtcccagaa acctgtggct gcttcaacca ttgacagttt  
61 tgctgctgct ggcttctgca gacagtcaag ctgcagctcc cccaaaggct gtgctgaaac  
121 ttgagccccc gtggatcaac gtgctccagg aggactctgt gactctgaca tgccaggggg  
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241 acacgcagcc cagctacagg ttcaaggcca acaacaatga cagcggggag tacacgtgcc  
301 agactggcca gaccagcctc agcgaccctg tgcattctgac tgtgctttcc gaatggctgg  
361 tgctccagac cctcacctg gagttccagg agggagaaac catcatgctg aggtgccaca  
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481 tctcccgttt ggatcccacc ttctccatcc cacaagcaaa ccacagtcac agtggtgatt  
541 accactgcac aggaacata ggctacacgc tgttctcacc caagcctgtg accatcactg  
601 tccaagtgcc cagcatgggc agctcttcac caatggggat cattgtggct gtggtcattg  
661 cgactgctgt agcagccatt gttgctgctg tagtggcctt gatctactgc aggaaaaagc  
721 ggatttcagc caattccact gatcctgtga aggctgcccc atttgagcca cctggacgtc  
781 aaatgattgc catcagaag agacaacttg aagaaaccaa caatgactat gaaacagctg  
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901 tgactcttcc tcccacgcac catgtcaaca gtaataacta aagagtaacg ttatgccatg  
961 tggtcatact ctacagcttg tgatggatga caaaaagagg ggaattgtta aaggaaaatt  
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1141 aagcaaaact taacttggat catttctggg aaatgcttat gttagaata agacaacccc  
1201 agccaatcac aagcagccta ctaacatata attaggtgac tagggacttt ctaagaagat  
1261 acctaccccc aaaaaacaa tatgtaattg aaaaccaacc gattgccttt attttgcttc  
1321 cacattttcc caataaatac ttgctgtga cattttgcca ctggaacact aaacttcagt  
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1801 cactgcttat gcagtcggaa gctccagaag aacaaagagc ccaattacca gaaccacatt  
1861 aagtctccat tgttttgcct tgggatttga gaagagaatt agagaggtga ggtctggta  
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2101 tctgttcac atccacacag ccaatacaat tagtcaaacc actgttatta acagatgtag  
2161 caacatgaga aacgcttatg ttacagggtta catgagagca atcatgtaag tctatatgac  
2221 ttcagaaatg ttaaaataga ctaacctcta acaacaaatt aaaagtgatt gtttcaagg  
2281 gatgcaatta ttgatgacct attttatttt tctataatga tcatatatta cctttgtaat  
2341 aaaacattat aacaaaaac

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## SEQ ID NO:17

human FcγRIII

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1      tctttggtga cttgtccact ccagtgtggc atcatgtggc agctgctcct cccaactgct
61     ctgctacttc tagtttcagc tggcatgcgg actgaagatc tcccaaaggc tgtggtgttc
121    ctggagcctc aatggtacag cgtgcttgag aaggacagtg tgactctgaa gtgccagggg
181    gcctactccc ctgaggacaa ttccacacag tggtttcaca atgagagcct catctcaagc
241    caggcctcga gctacttcat tgacgtgcc acagtcaacg acagtggaga gtacagggtc
301    cagacaaacc tctccacct cagtgacctg gtgcagctag aagtccatat cggctggctg
361    ttgctccagg cccctcggtg ggtgttcaag gaggaagacc ctattcacct gaggtgtcac
421    agctggaaga aactgctct gcataaggtc acatatttac agaatggcaa agacaggaag
481    tattttcac ataattctga cttccacatt ccaaaagcca cactcaaaga tagcggctcc
541    tacttctgca gggggcttgt tgggagtaaa aatgtgtctt cagagactgt gaacatcacc
601    atcactcaag gtttggcagt gtcaaccatc tcattcattct ctccacctgg gtaccaagtc
661    tctttctgct tggatggtt actccttttt gcagtggaca caggactata tttctctgtg
721    aagacaaaca tttgaagctc aacaagagac tggaaggacc ataaacttaa atggagaaaag
781    gaccctcaag acaaatgacc cccatcccat gggagtaata agagcagtgg cagcagcatc
841    tctgaacatt tctctggatt tgcaacccca tcctcctcag gcctctc
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SEQ ID NO: 18

## human FCERII

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accccaacac  
61 actaggagga cagacacagg ctccaaactc cactaagtga ccagagctgt gattgtgccc  
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241 ggcgtgggac tcagatcgtg ctgctggggc tggtagccgc cgctctgtgg gctgggctgc  
301 tgactctgct tctcctgtgg cactgggaca ccacacagag tctaaaacag ctggaagaga  
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421 tggcgagaa atcccagtcc acgcagattt cacaggaact ggaggaactt cgagctgaac  
481 agcagagatt gaaatctcag gacttgagc tgcctggaa cctgaacggg cttcaagcag  
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